

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Burton, Dennis R.
Burioni, Roberto
Williamson, R. Anthony
Sanna, Pietro Paolo

(ii) TITLE OF INVENTION: LIGAND CAPTURE-DIRECTED SELECTION OF ANTIBODY

(iii) NUMBER OF SEQUENCES: 23

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Spensley Horn Jubas & Lubitz
(B) STREET: 1880 Century Park East, Suite 500
(C) CITY: Los Angeles
(D) STATE: California
(E) COUNTRY: USA
(F) ZIP: 90067

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US
(B) FILING DATE: October 3, 1994
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: LISA A. HAILE, PH.D.
(B) REGISTRATION NUMBER: P-38,347
(C) REFERENCE/DOCKET NUMBER: PD-3809

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (619) 455-5100
(B) TELEFAX: (619) 455-5110

(2) INFORMATION FOR SEQ ID NO:1

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 32 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N1

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..32

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1

GGCCGCAAAT TCTATTTCAA GGAGACAGTC AT

32

(2) INFORMATION FOR SEQ ID NO:2

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 36 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N2

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..36

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

AATGAAATAC CTATTGCCTA CGGCAGCCGC TGGATT

36

10079802-021400

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 32 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:
 (B) CLONE: N3

(ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..32

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

GTTATTACTC GCTGCCCAAC CAGCCATGGC CC

32

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 29 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:
 (B) CLONE: N6

(ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..29

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CAGTTTCACC TGGGCCATGG CTGGTTGGG

29

10079300-041901

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 40 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N7

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

CAGCGAGTAA TAACAATCCA GCGGCTGCCG TAGGCAATAG

40

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 38 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N8

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..38

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

GTATTCATT ATGACTGTCT CCTTGAAATA GAATTTGC

38

10079602-071906

(2) INFORMATION FOR SEQ ID NO:7:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 40 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N9-4

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

AGGTGAAACT GCTCGAGATT TCTAGACTAG TTACCCGTAC

40

(2) INFORMATION FOR SEQ ID NO:8 :

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 38 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N10-5

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..38

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8 :

CGGAACGTCG TACGGGTAAC TAGTCTAGAA ATCTCGAG

38

10070803-021902

(2) INFORMATION FOR SEQ ID NO:9 :

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 33 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N11

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..33

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9 :

GACGTTCCGG ACTACGGTTC TTAATAGAAT TCG

33

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

- (B) CLONE: N12

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..28

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

TCGACGAATT CTATTAAGAA CCGTAGTC

28

205720"20364001

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 34 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..34

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

TGAATTCTAA ACTAGTCGCC AAGGAGACAG TCAT

34

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 36 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..36

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

AATGAAATAC CTATTGCCTA CGGCAGCCGC TGGATT

36

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 31 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..31

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GTTATTACTC GCTGCCCAAC CAGCCATGGC C

31

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 30 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..30

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

GAGCTCGTCA GTTCTAGAGT TAAGCGGCCG

30

(2) INFORMATION FOR SEQ ID NO:15:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 48 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..48

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

5 CTATTTTCATT ATGACTGTCT CCTTGGCGAC TAGTTTAGAA TTCAAGCT 48

(2) INFORMATION FOR SEQ ID NO:16:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 40 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

CAGCGAGTAA TAACAATCCA GCGGCTGCCG TAGGCAATAG 40

(2) INFORMATION FOR SEQ ID NO:17:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 27 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..27

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

TGACGAGCTC GGCCATGGCT GGTGGG

27

(2) INFORMATION FOR SEQ ID NO:18:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..24

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

TCGACGGCCG CTTAACTCTA GAAC

24

(2) INFORMATION FOR SEQ ID NO:19:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 48 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..48

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

GAGACGACTA GTGGTGGCGG TGGCTCTCCA TTCGTTTGTG AATATCAA

48

(2) INFORMATION FOR SEQ ID NO:20:

10079607-024902
205120-20867007

(D) TOPOLOGY: linear

(B) LOCATION: 1..40

(D) TOPOLOGY: linear

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..27

5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

ACCGAGCTCG AATTCGTAAT CATGGTC

27

(2) INFORMATION FOR SEQ ID NO:23:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..31

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

AGCTGTTGAA TTCGTGAAAT TGTTATCCGC T

31

10079301-031903

The foregoing is meant to illustrate, but not to limit, the scope of the invention. Indeed, those of ordinary skill in the art can readily envision and produce further embodiments, based on the teachings herein, without undue experimentation.

10079802, 021902

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Burton, Dennis R.
Burioni, Roberto
Williamson, R. Anthony
Sanna, Pietro Paolo
- (ii) TITLE OF INVENTION: LIGAND CAPTURE-DIRECTED SELECTION OF ANTIBODY
- (iii) NUMBER OF SEQUENCES: 23
- (iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Spensley Horn Jubas & Lubitz
(B) STREET: 1880 Century Park East, Suite 500
(C) CITY: Los Angeles
(D) STATE: California
(E) COUNTRY: USA
(F) ZIP: 90067
- (v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER: US
(B) FILING DATE: October 3, 1994
(C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: LISA A. HAILE, PH.D.
(B) REGISTRATION NUMBER: P-38,347
(C) REFERENCE/DOCKET NUMBER: PD-3809
- (ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: (619) 455-5100
(B) TELEFAX: (619) 455-5110

(2) INFORMATION FOR SEQ ID NO:1

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 32 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (vii) IMMEDIATE SOURCE:
(B) CLONE: N1
- (ix) FEATURE:
(A) NAME/KEY: CDS
(B) LOCATION: 1..32
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1

32

[illegible]

- (ii) MOLECULE TYPE: DNA (genomic)

- ```
(ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..36
```

- AATGAAATAC CTATTGCCTA CGGCAGCCGC TGGATT

36

## (ii) MOLECULE TYPE: DNA (genomic)

- (ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..32

- GTTATTACTC GCTGCCCAAC CAGCCATGGC CC

32

## (ii) MOLECULE TYPE: DNA (genomic)

- (ix) FEATURE:  
(A) NAME/KEY: CDS

(B) LOCATION: 1..29

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CAGTTTCACC TGGGCCATGG CTGGTTGGG

29

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 40 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

(B) CLONE: N7

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

CAGCGAGTAA TAACAATCCA GCGGCTGCCG TAGGCAATAG

40

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 38 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:

(B) CLONE: N8

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..38

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

GTATTTTCATT ATGACTGTCT CCTTGAAATA GAATTTGC

38

(2) INFORMATION FOR SEQ ID NO:7:

100706072006000



- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 40 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:  
 (B) CLONE: N9-4

(ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

AGGTGAAACT GCTCGAGATT TCTAGACTAG TTACCCGTAC

40

(2) INFORMATION FOR SEQ ID NO:8 :

- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 38 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:  
 (B) CLONE: N10-5

(ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..38

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8 :

CGGAACGTCG TACGGGTAAC TAGTCTAGAA ATCTCGAG

38

(2) INFORMATION FOR SEQ ID NO:9 :

- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 33 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(vii) IMMEDIATE SOURCE:  
 (B) CLONE: N11

(ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..33

10079800-021900

| Variable       | Mean | Standard Deviation | Minimum | Maximum |
|----------------|------|--------------------|---------|---------|
| Age            | 34.5 | 10.2               | 22      | 55      |
| Gender         | 0.5  | 0.5                | 0       | 1       |
| Marital Status | 0.6  | 0.5                | 0       | 1       |
| Education      | 12.5 | 1.5                | 10      | 16      |
| Income         | 3500 | 1500               | 1000    | 8000    |
| Health         | 0.8  | 0.2                | 0       | 1       |
| Smoking        | 0.3  | 0.5                | 0       | 1       |
| Alcohol        | 0.2  | 0.4                | 0       | 1       |
| Exercise       | 0.4  | 0.5                | 0       | 1       |
| Stress         | 0.6  | 0.5                | 0       | 1       |
| Sleep          | 0.7  | 0.3                | 0       | 1       |
| Diet           | 0.5  | 0.5                | 0       | 1       |
| Work           | 0.8  | 0.2                | 0       | 1       |
| Family         | 0.6  | 0.5                | 0       | 1       |
| Friends        | 0.7  | 0.4                | 0       | 1       |
| Hobbies        | 0.5  | 0.5                | 0       | 1       |
| Travel         | 0.4  | 0.5                | 0       | 1       |
| Shopping       | 0.6  | 0.5                | 0       | 1       |
| Reading        | 0.5  | 0.5                | 0       | 1       |
| TV             | 0.7  | 0.4                | 0       | 1       |
| Music          | 0.6  | 0.5                | 0       | 1       |
| Gardening      | 0.4  | 0.5                | 0       | 1       |
| Cooking        | 0.5  | 0.5                | 0       | 1       |
| Volunteering   | 0.3  | 0.5                | 0       | 1       |
| Religion       | 0.5  | 0.5                | 0       | 1       |
| Politics       | 0.4  | 0.5                | 0       | 1       |
| Environment    | 0.6  | 0.5                | 0       | 1       |
| Technology     | 0.7  | 0.4                | 0       | 1       |
| Art            | 0.5  | 0.5                | 0       | 1       |
| Sports         | 0.6  | 0.5                | 0       | 1       |
| Traveling      | 0.4  | 0.5                | 0       | 1       |
| Shopping       | 0.5  | 0.5                | 0       | 1       |
| Reading        | 0.6  | 0.5                | 0       | 1       |
| TV             | 0.7  | 0.4                | 0       | 1       |
| Music          | 0.6  | 0.5                | 0       | 1       |
| Gardening      | 0.4  | 0.5                | 0       | 1       |
| Cooking        | 0.5  | 0.5                | 0       | 1       |
| Volunteering   | 0.3  | 0.5                | 0       | 1       |
| Religion       | 0.5  | 0.5                | 0       | 1       |
| Politics       | 0.4  | 0.5                | 0       | 1       |
| Environment    | 0.6  | 0.5                | 0       | 1       |
| Technology     | 0.7  | 0.4                | 0       | 1       |
| Art            | 0.5  | 0.5                | 0       | 1       |
| Sports         | 0.6  | 0.5                | 0       | 1       |
| Traveling      | 0.4  | 0.5                | 0       | 1       |
| Shopping       | 0.5  | 0.5                | 0       | 1       |
| Reading        | 0.6  | 0.5                | 0       | 1       |
| TV             | 0.7  | 0.4                | 0       | 1       |
| Music          | 0.6  | 0.5                | 0       | 1       |
| Gardening      | 0.4  | 0.5                | 0       | 1       |
| Cooking        | 0.5  | 0.5                | 0       | 1       |
| Volunteering   | 0.3  | 0.5                | 0       | 1       |
| Religion       | 0.5  | 0.5                | 0       | 1       |
| Politics       | 0.4  | 0.5                | 0       | 1       |
| Environment    | 0.6  | 0.5                | 0       | 1       |
| Technology     | 0.7  | 0.4                | 0       | 1       |
| Art            | 0.5  | 0.5                | 0       | 1       |
| Sports         | 0.6  | 0.5                | 0       | 1       |
| Traveling      | 0.4  | 0.5                | 0       | 1       |
| Shopping       | 0.5  | 0.5                | 0       | 1       |
| Reading        | 0.6  | 0.5                | 0       | 1       |
| TV             | 0.7  | 0.4                | 0       | 1       |
| Music          | 0.6  | 0.5                | 0       | 1       |
| Gardening      | 0.4  | 0.5                | 0       | 1       |
| Cooking        | 0.5  | 0.5                | 0       | 1       |
| Volunteering   | 0.3  | 0.5                | 0       | 1       |
| Religion       | 0.5  | 0.5                | 0       | 1       |
| Politics       | 0.4  | 0.5                | 0       | 1       |
| Environment    | 0.6  | 0.5                | 0       | 1       |
| Technology     | 0.7  | 0.4                | 0       | 1       |
| Art            | 0.5  | 0.5                | 0       | 1       |
| Sports         | 0.6  | 0.5                | 0       | 1       |
| Traveling      | 0.4  | 0.5                | 0       | 1       |
| Shopping       | 0.5  | 0.5                | 0       | 1       |
| Reading        | 0.6  | 0.5                | 0       | 1       |
| TV             | 0.7  | 0.4                | 0       | 1       |
| Music          | 0.6  | 0.5                | 0       | 1       |
| Gardening      | 0.4  | 0.5                | 0       | 1       |
| Cooking        | 0.5  | 0.5                | 0       | 1       |
| Volunteering   | 0.3  | 0.5                | 0       | 1       |
| Religion       | 0.5  | 0.5                | 0       | 1       |
| Politics       | 0.4  | 0.5                | 0       | 1       |
| Environment    | 0.6  | 0.5                | 0       | 1       |
| Technology     | 0.7  | 0.4                | 0       | 1       |
| Art            | 0.5  | 0.5                | 0       | 1       |
| Sports         | 0.6  | 0.5                | 0       | 1       |
| Traveling      | 0.4  | 0.5                |         |         |

33

(2) INFORMATION FOR SEQ ID NO:10:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 28 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (vii) IMMEDIATE SOURCE:
  - (B) CLONE: N12

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

28

(2) INFORMATION FOR SEQ ID NO:11:

- ```
(i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 34 base pairs
      (B) TYPE: nucleic acid
      (C) STRANDEDNESS: single
      (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:
      (A) NAME/KEY: CDS
      (B) LOCATION: 1..34
```

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

34

(2) INFORMATION FOR SEQ ID NO:12:

- ```
(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 36 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..36
```

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

AATGAAATAC CTATTGCCTA CGGCAGCCGC TGGATT

36

(2) INFORMATION FOR SEQ ID NO:13:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 31 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..31

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GTTATTACTC GCTGCCCAAC CAGCCATGGC C

31

(2) INFORMATION FOR SEQ ID NO:14:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 30 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..30

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

GAGCTCGTCA GTTCTAGAGT TAAGCGGCCG

30

(2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 48 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..48

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

CTATTTTATT ATGACTGTCT CCTTGGCGAC TAGTTTAGAA TTCAAGCT

48

(2) INFORMATION FOR SEQ ID NO:16:

20061201 20061201

- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 40 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

- (ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

CAGCGAGTAA TAACAATCCA GCGGCTGCCG TAGGCAATAG

40

(2) INFORMATION FOR SEQ ID NO:17:

- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 27 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

- (ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..27

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

TGACGAGCTC GGCCATGGCT GGTG

27

(2) INFORMATION FOR SEQ ID NO:18:

- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 24 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

- (ix) FEATURE:  
 (A) NAME/KEY: CDS  
 (B) LOCATION: 1..24

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

TCGACGGCCG CTTAACTCTA GAAC

24

(2) INFORMATION FOR SEQ ID NO:19:

- (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 48 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..48

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

GAGACGACTA GTGGTGGCGG TGGCTCTCCA TTCGTTTGTG AATATCAA

48

(2) INFORMATION FOR SEQ ID NO:20:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 40 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

TTACTAGCTA GCATAATAAC GGAATACCCA AAAGAAGTGG

40

(2) INFORMATION FOR SEQ ID NO:21:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 36 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 1..36

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

TATGCTAGCT AGTAACACGA CAGGTTTCCC GACTGG

36

(2) INFORMATION FOR SEQ ID NO:22:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

20250720 2025/07/20

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..27

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

ACCGAGCTCG AATTCGTAAT CATGGTC

27

(2) INFORMATION FOR SEQ ID NO:23:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 31 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..31

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

AGCTGTTGAA TTCGTGAAAT TGTTATCCGC T

31

2025-07-20 20:20:00